

(SEQUENCE LISTING)

SEQ ID NO: 1

Length of sequence: 18

Type of sequence: Nucleic acid

5 Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +176 to +193 of

10 SEQ ID NO:9

Sequence:

AGAAAGTCTT CACTCTGC

SEQ ID NO: 2

15 Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

20 Antisense: YES

Feature of sequence: Corresponding to from +181 to +198 of

SEQ ID NO:9

Sequence:

TTGAAAGAAA GTCTTCAC

25

SEQ ID NO: 3

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

5 Antisense: YES

Feature of sequence: Corresponding to from +367 to +384 of
SEQ ID NO:9

Sequence:

GGTCTTCAGG TTCTCCCC

10

SEQ ID NO: 4

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

15 Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +637 to +654 of
SEQ ID NO:9

20 Sequence:

CTGGGTCAGC TATCCCAG

SEQ ID NO: 5

Length of sequence: 18

25 Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +915 to +932 of
SEQ ID NO:9

5 Sequence:

GCTTGAATG GAAGCTTC

SEQ ID NO: 6

Length of sequence: 18

10 Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Synthesized DNA

Antisense: YES

15 Feature of sequence: Corresponding to from +1246 to +1263 of
SEQ ID NO:9

Sequence:

GGCTGGTTAG GAACTCCT

20 SEQ ID NO: 7

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

25 Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +1249 to +1266 of

SEQ ID NO:9

Sequence:

CCAGGCTGGT TAGGAACT

5 SEQ ID NO: 8

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

10 Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Mouse IL-10 protein gene

Sequence:

AGGTCCTGGA GTCCAGCA

15

SEQ ID NO: 9

Length of sequence: 1601

Type of sequence: Nucleic acid

Number of strand: Single-strand

20 Topology: Straight

Kind of sequence: cDNA

Antisense: NO

Feature of sequence: cDNA of Human IL-10 protein

Sequence:

25 AAACCACAAG ACAGACTTGC AAAAGAAGGC ATGCACAGCT CAGCACTGCT CTGTTGCCTG 60
GTCCTCCTGA CTGGGGTGAG GGCCAGCCCA GGCCAGGGCA CCCAGTCTGA GAACAGCTGC 120
ACCCACTTCC CAGGCAACCT GCCTAACATG CTTGAGATC TCCGAGATGC CTTCAGCAGA 180

GTGAAGACTT TCTTTCAAAT GAAGGATCAG CTGGACAAC TGTGTTAAA GGAGTCCTTG 240
CTGGAGGACT TTAAGGGTTA CCTGGGTTGC CAAGCCTTGT CTGAGATGAT CCAGTTTTAC 300
CTGGAGGAGG TGATGCCCCA AGCTGAGAAC CAAGACCCAG ACATCAAGGC GCATGTGAAC 360
TCCCTGGGGG AGAACCTGAA GACCCTCAGG CTGAGGCTAC GCGCTGTCA TCGATTTCTT 420
5 CCCTGTGAAA ACAAGAGCAA GGCCGTGGAG CAGGTGAAGA ATGCCTTTAA TAAGCTCCAA 480
GAGAAAGGCA TCTACAAAGC CATGAGTGAG TTTGACATCT TCATCAACTA CATAGAAGCC 540
TACATGACAA TGAAGATACG AACTGAGAC ATCAGGGTGG CGACTCTATA GACTCTAGGA 600
CATAAATTAG AGGTCTCCAA AATCGGATCT GGGGCTCTGG GATAGCTGAC CCAGCCCCTT 660
GAGAAACCTT ATTGTACCTC TCTTATAGAA TATTTATTAC CTCTGATACC TCAACCCCCA 720
10 TTTCTATTTA TTTACTGAGC TTCTCTGTGA ACGATTTAGA AAGAAGCCCA ATATTATAAT 780
TTTTTTCAAT ATTTATTATT TTCACCTGTT TTTAAGCTGT TTCCATAGGG TGACACACTA 840
TGGTATTTGA GTGTTTTAAG ATAAATTATA AGTTACATAA GGGAGGAAAA AAAATGTTCT 900
TTGGGGAGCC AACAGAAGCT TCCATTCCAA GCCTGACCAC GCTTTCTAGC TGTTGAGCTG 960
TTTTCCCTGA CCTCCCTCTA ATTTATCTTG TCTCTGGGCT TGGGGCTTCC TAACTGCTAC 1020
15 AAATACTCTT AGGAAGAGAA ACCAGGGAGC CCCTTGATG ATTAATTCAC CTTCAGTGT 1080
CTCGGAGGGA TTCCCCTAAC CTCATTCCCC AACCATTCA TTCTTGAAAG CTGTGGCCAG 1140
CTTGTTATTT ATAACAACCT AAATTTGGTT CTAGGCCGGG CGCGGTGGCT CACGCCTGTA 1200
ATCCCAGCAC TTTGGGAGGC TGAGGCGGGT GGATCACTTG AGGTCAGGAG TTCCTAACCA 1260
GCCTGGTCAA CATGGTGAAA CCCCCTCTCT ACTAAAAATA CAAAAATTAG CCGGGCATGG 1320
20 TGGCGCGCAC CTGTAATCCC AGCTACTTGG GAGGCTGAGG CAAGAGAATT GCTTGAACCC 1380
AGGAGATGGA AGTTGCAGTG AGCTGATATC ATGCCCCTGT ACTCCAGCCT GGGTGACAGA 1440
GCAAGACTCT GTCTCAAAAA AATAAAAAATA AAAATAAATT TGTTTCTAAT AGAACTCAGT 1500
TTTAACTAGA ATTTATTCAA TTCCTCTGGG AATGTTACAT TGTTTGTCTG TCTTCATAGC 1560
AGATTTTAAT TTTGAATAAA TAAATGTATC TTATTACAT C 1601